

#7

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/511,722

Source: PT/10

Date Processed by STIC: 8/4/05

ENTERED



PCT

RAW SEQUENCE LISTING

DATE: 08/04/2005

PATENT APPLICATION: US/10/511,722

TIME: 14:19:01

Input Set : A:\sequence listing.txt

Output Set: N:\CRF4\08042005\J511722.raw

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3 <110> APPLICANT: Yeda Research and Development Co. Ltd.
4     Wallach, David
5     Shmushkovich , Taisia
6     Ramakrishnan, Parameswaran
8 <120> TITLE OF INVENTION: Derivatives of the IL-2 receptor Gamma chain, their
preparation and use
10 <130> FILE REFERENCE: 530
C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/511,722
C--> 12 <141> CURRENT FILING DATE: 2004-10-18
12 <150> PRIOR APPLICATION NUMBER: 149217
13 <151> PRIOR FILING DATE: 2002-04-18
15 <150> PRIOR APPLICATION NUMBER: 152183
16 <151> PRIOR FILING DATE: 2002-10-08
18 <160> NUMBER OF SEQ ID NOS: 21
20 <170> SOFTWARE: PatentIn version 3.1
22 <210> SEQ ID NO: 1
23 <211> LENGTH: 86
24 <212> TYPE: PRT
25 <213> ORGANISM: Homo sapiens
27 <400> SEQUENCE: 1
29 Glu Arg Thr Met Pro Arg Ile Pro Thr Leu Lys Asn Leu Glu Asp Leu
30 1           5           10           15
33 Val Thr Glu Tyr His Gly Asn Phe Ser Ala Trp Ser Gly Val Ser Lys
34           20           25           30
37 Gly Leu Ala Glu Ser Leu Gln Pro Asp Tyr Ser Glu Arg Leu Cys Leu
38           35           40           45
41 Val Ser Glu Ile Pro Pro Lys Gly Gly Ala Leu Gly Glu Gly Pro Gly
42           50           55           60
45 Ala Ser Pro Cys Asn Gln His Ser Pro Tyr Trp Ala Pro Pro Cys Tyr
46 65           70           75           80
49 Thr Leu Lys Pro Glu Thr
50           85
53 <210> SEQ ID NO: 2
54 <211> LENGTH: 41
55 <212> TYPE: PRT
56 <213> ORGANISM: Homo sapiens
58 <400> SEQUENCE: 2
60 Leu Cys Leu Val Ser Glu Ile Pro Pro Lys Gly Gly Ala Leu Gly Glu
61 1           5           10           15
64 Gly Pro Gly Ala Ser Pro Cys Asn Gln His Ser Pro Tyr Trp Ala Pro
65           20           25           30
68 Pro Cys Tyr Thr Leu Lys Pro Glu Thr
69           35           40
72 <210> SEQ ID NO: 3

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73 <211> LENGTH: 12
74 <212> TYPE: PRT
75 <213> ORGANISM: Homo sapiens
77 <400> SEQUENCE: 3
79 Trp Ala Pro Pro Cys Tyr Thr Leu Lys Pro Glu Thr
80 1          5          10
83 <210> SEQ ID NO: 4
84 <211> LENGTH: 39
85 <212> TYPE: DNA
86 <213> ORGANISM: Homo sapiens
88 <400> SEQUENCE: 4
89 tgggcccccc catgttacac cctaaagcct gaaacctga          39
92 <210> SEQ ID NO: 5
93 <211> LENGTH: 261
94 <212> TYPE: DNA
95 <213> ORGANISM: Homo sapiens
97 <400> SEQUENCE: 5
98 gaacggacga tgccccgaat tcccaccctg aagaacctag aggatcttgt tactgaatac          60
100 cacgggaact tttcggcctg gagtgggtgtg tctaagggaac tggtgagag tctgcagcca          120
102 gactacagtg aacgactctg cctcgtcagt gagattcccc caaaaggagg ggccttggg          180
104 gaggggcctg gggcctcccc atgcaaccag catagcccct actgggcccc cccatgttac          240
106 accctaaagc ctgaaacctg a          261
109 <210> SEQ ID NO: 6
110 <211> LENGTH: 126
111 <212> TYPE: DNA
112 <213> ORGANISM: Homo sapiens
114 <400> SEQUENCE: 6
115 ctctgcctcg tcaagtgaat tccccaaaaa ggaggggccc ttggggaggg gcctggggcc          60
117 tccccatgca accagcatag cccctactgg gcccccccat gttacacct aaagcctgaa          120
119 acctga          126
122 <210> SEQ ID NO: 7
123 <211> LENGTH: 37
124 <212> TYPE: DNA
125 <213> ORGANISM: Homo sapiens
127 <400> SEQUENCE: 7
128 ctcgtcagt agattgccgc aaaaggagg ggccttg          37
131 <210> SEQ ID NO: 8
132 <211> LENGTH: 37
133 <212> TYPE: DNA
134 <213> ORGANISM: Homo sapiens
136 <400> SEQUENCE: 8
137 caagggccccc tctttttgcg gcaatctcac tgacgag          37
140 <210> SEQ ID NO: 9
141 <211> LENGTH: 35
142 <212> TYPE: DNA
143 <213> ORGANISM: Homo sapiens
145 <400> SEQUENCE: 9
146 gccctactg ggccgccgca tgttacaccc taaag          35
149 <210> SEQ ID NO: 10

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150 <211> LENGTH: 35
151 <212> TYPE: DNA
152 <213> ORGANISM: Homo sapiens
154 <400> SEQUENCE: 10
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158 <210> SEQ ID NO: 11
159 <211> LENGTH: 39
160 <212> TYPE: DNA
161 <213> ORGANISM: Homo sapiens
163 <400> SEQUENCE: 11
164 gtcagtgaga ttccccagc aggaggggcc cttggggag 39
167 <210> SEQ ID NO: 12
168 <211> LENGTH: 39
169 <212> TYPE: DNA
170 <213> ORGANISM: Homo sapiens
172 <400> SEQUENCE: 12
173 ctccccaagg gccctcctg ctgggggaat ctactgac 39
176 <210> SEQ ID NO: 13
177 <211> LENGTH: 33
178 <212> TYPE: DNA
179 <213> ORGANISM: Homo sapiens
181 <400> SEQUENCE: 13
182 ggaggggccc ttggggcggg gcctggggcc tcc 33
185 <210> SEQ ID NO: 14
186 <211> LENGTH: 33
187 <212> TYPE: DNA
188 <213> ORGANISM: Homo sapiens
190 <400> SEQUENCE: 14
191 ggaggcccca ggccccgccc caagggcccc tcc 33
194 <210> SEQ ID NO: 15
195 <211> LENGTH: 33
196 <212> TYPE: DNA
197 <213> ORGANISM: Homo sapiens
199 <400> SEQUENCE: 15
200 cagcatagcc cctacgcggc cccccatgt tac 33
203 <210> SEQ ID NO: 16
204 <211> LENGTH: 33
205 <212> TYPE: DNA
206 <213> ORGANISM: HOMO SAPIENS
208 <400> SEQUENCE: 16
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212 <210> SEQ ID NO: 17
213 <211> LENGTH: 44
214 <212> TYPE: PRT
215 <213> ORGANISM: Homo sapiens
217 <400> SEQUENCE: 17
219 Trp Leu Glu Arg Thr Met Pro Arg Ile Pro Thr Leu Lys Asn Leu Glu
220 1 5 10 15
223 Asp Leu Val Thr Glu Tyr His Gly Asn Phe Ser Ala Trp Ser Gly Val

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224          20          25          30
227 Ser Lys Gly Leu Ala Glu Ser Leu Gln Pro Asp Tyr
228          35          40
231 <210> SEQ ID NO: 18
232 <211> LENGTH: 81
233 <212> TYPE: PRT
234 <213> ORGANISM: Homo sapiens
236 <400> SEQUENCE: 18
238 His Arg Val Ser Ala Ala Glu Leu Gly Gly Lys Val Asn Arg Ala Leu
239 1          5          10          15
242 Gln Gln Val Gly Gly Leu Lys Ser Pro Trp Arg Gly Glu Tyr Lys Glu
243          20          25          30
246 Pro Arg His Pro Pro Pro Asn Gln Ala Asn Tyr His Gln Thr Leu His
247          35          40          45
250 Ala Gln Pro Arg Glu Leu Ser Pro Arg Ala Pro Gly Pro Arg Pro Ala
251          50          55          60
254 Glu Glu Thr Thr Gly Arg Ala Pro Lys Leu Gln Pro Pro Leu Pro Pro
255 65          70          75          80
258 Glu
262 <210> SEQ ID NO: 19
263 <211> LENGTH: 324
264 <212> TYPE: PRT
265 <213> ORGANISM: Homo sapiens
267 <400> SEQUENCE: 19
269 Pro Leu Thr Ala Gln Ala Ile Gln Glu Gly Leu Arg Lys Glu Pro Ile
270 1          5          10          15
273 His Arg Val Ser Ala Ala Glu Leu Gly Gly Lys Val Asn Arg Ala Leu
274          20          25          30
277 Gln Gln Val Gly Gly Leu Lys Ser Pro Trp Arg Gly Glu Tyr Lys Glu
278          35          40          45
281 Pro Arg His Pro Pro Pro Asn Gln Ala Asn Tyr His Gln Thr Leu His
282          50          55          60
285 Ala Gln Pro Arg Glu Leu Ser Pro Arg Ala Pro Gly Pro Arg Pro Ala
286 65          70          75          80
289 Glu Glu Thr Thr Gly Arg Ala Pro Lys Leu Gln Pro Pro Leu Pro Pro
290          85          90          95
293 Glu Pro Pro Glu Pro Asn Lys Ser Pro Pro Leu Thr Leu Ser Lys Glu
294          100         105         110
297 Glu Ser Gly Met Trp Glu Pro Leu Pro Leu Ser Ser Leu Glu Pro Ala
298          115         120         125
301 Pro Ala Arg Asn Pro Ser Ser Pro Glu Arg Lys Ala Thr Val Pro Glu
302          130         135         140
305 Gln Glu Leu Gln Gln Leu Glu Ile Glu Leu Phe Leu Asn Ser Leu Ser
306 145         150         155         160
309 Gln Pro Phe Ser Leu Glu Glu Gln Glu Gln Ile Leu Ser Cys Leu Ser
310          165         170         175
313 Ile Asp Ser Leu Ser Leu Ser Asp Asp Ser Glu Lys Asn Pro Ser Lys
314          180         185         190
317 Ala Ser Gln Ser Ser Arg Asp Thr Leu Ser Ser Gly Val His Ser Trp

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318      195      200      205
321 Ser Ser Gln Ala Glu Ala Arg Ser Ser Ser Trp Asn Met Val Leu Ala
322      210      215      220
325 Arg Gly Arg Pro Thr Asp Thr Pro Ser Tyr Phe Asn Gly Val Lys Val
326 225      230      235      240
329 Gln Ile Gln Ser Leu Asn Gly Glu His Leu His Ile Arg Glu Phe His
330      245      250      255
333 Arg Val Lys Val Gly Asp Ile Ala Thr Gly Ile Ser Ser Gln Ile Pro
334      260      265      270
337 Ala Ala Ala Phe Ser Leu Val Thr Lys Asp Gly Gln Pro Val Arg Tyr
338      275      280      285
341 Asp Met Glu Val Pro Asp Ser Gly Ile Asp Leu Gln Cys Thr Leu Ala
342      290      295      300
345 Pro Asp Gly Ser Phe Ala Trp Ser Trp Arg Val Lys His Gly Gln Leu
346 305      310      315      320
349 Glu Asn Arg Pro
353 <210> SEQ ID NO: 20
354 <211> LENGTH: 357
355 <212> TYPE: PRT
356 <213> ORGANISM: Homo sapiens
358 <400> SEQUENCE: 20
360 Met Leu Lys Pro Ser Leu Pro Phe Thr Ser Leu Leu Phe Leu Gln Leu
361 1      5      10      15
364 Pro Leu Leu Gly Val Gly Leu Asn Thr Thr Ile Leu Thr Pro Asn Gly
365      20      25      30
368 Asn Glu Asp Thr Thr Ala Asp Phe Phe Leu Thr Thr Met Pro Thr Asp
369      35      40      45
372 Ser Leu Ser Val Ser Thr Leu Pro Leu Pro Glu Val Gln Cys Phe Val
373      50      55      60
376 Phe Asn Val Glu Tyr Met Asn Cys Thr Trp Asn Ser Ser Ser Glu Pro
377 65      70      75      80
380 Gln Pro Thr Asn Leu Thr Leu His Tyr Trp Tyr Lys Asn Ser Asp Asn
381      85      90      95
384 Asp Lys Val Gln Lys Cys Ser His Tyr Leu Phe Ser Glu Glu Ile Thr
385      100      105      110
388 Ser Gly Cys Gln Leu Gln Lys Lys Glu Ile His Leu Tyr Gln Thr Phe
389      115      120      125
392 Val Val Gln Leu Gln Asp Pro Arg Glu Pro Arg Arg Gln Ala Thr Gln
393      130      135      140
396 Met Leu Lys Leu Gln Asn Leu Val Ile Pro Trp Ala Pro Glu Asn Leu
397 145      150      155      160
400 Thr Leu His Lys Leu Ser Glu Ser Gln Leu Glu Leu Asn Trp Asn Asn
401      165      170      175
404 Arg Phe Leu Asn His Cys Leu Glu His Leu Val Gln Tyr Arg Thr Asp
405      180      185      190
408 Trp Asp His Ser Trp Thr Glu Gln Ser Val Asp Tyr Arg His Lys Phe
409      195      200      205
412 Ser Leu Pro Ser Val Asp Gly Gln Lys Arg Tyr Thr Phe Arg Val Arg
413      210      215      220

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/511,722

DATE: 08/04/2005
TIME: 14:19:02

Input Set : A:\sequence listing.txt
Output Set: N:\CRF4\08042005\J511722.raw

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 8

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/511,722

DATE: 08/04/2005

TIME: 14:19:02

Input Set : A:\sequence listing.txt

Output Set: N:\CRF4\08042005\J511722.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No

L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date